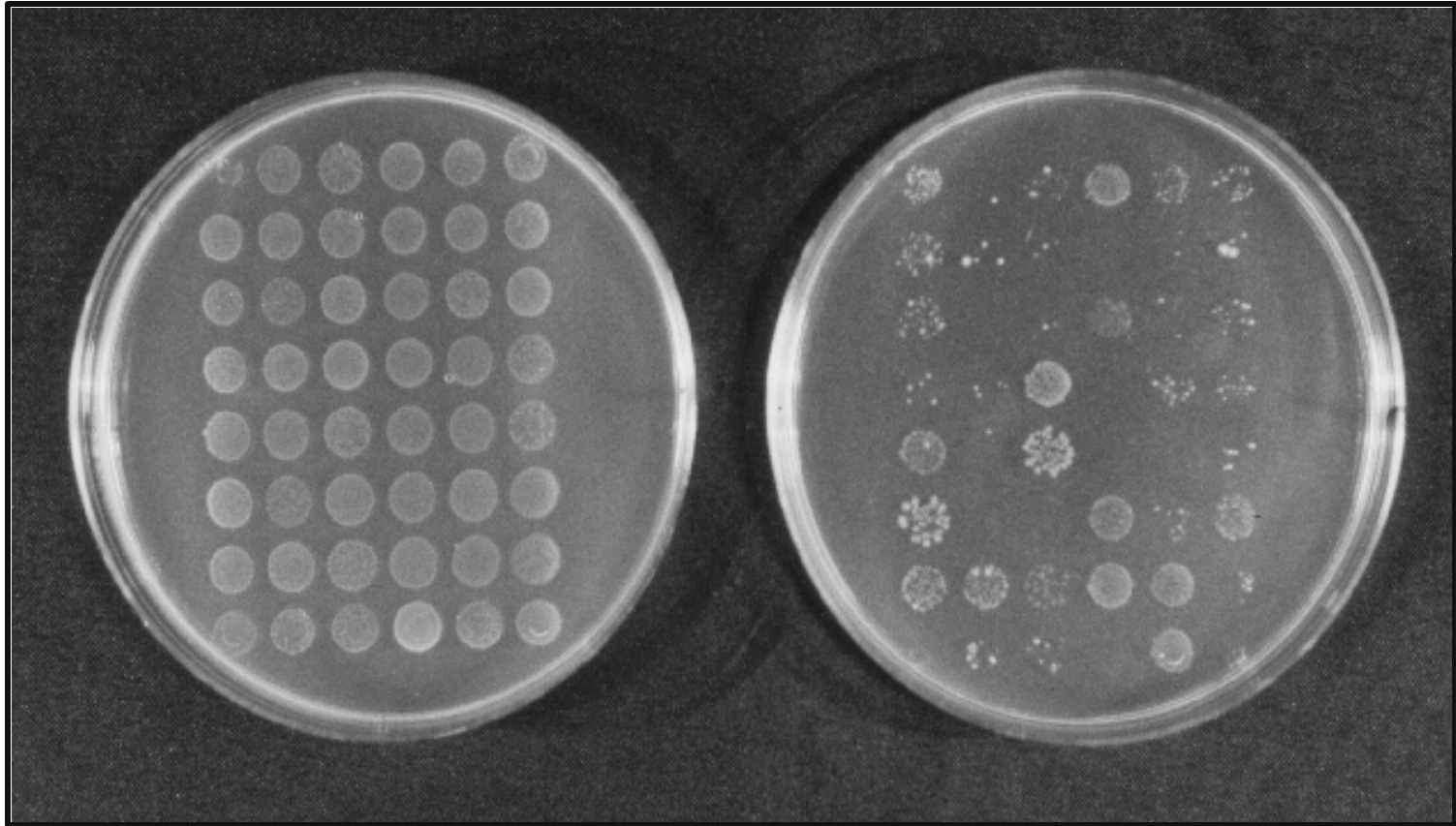


Microbial Source Tracking

- Phenotypic methods (biochemical based)
 - Antibiotic resistance analysis (ARA)
 - Carbon utilization profile (CUP)
- * Both are library dependent methods

Antibiotic Resistance Analysis



Patterns of Antibiotic Resistance

Antibiotic (mg/l)	Poultry	Humans
----% of isolate resistant----		
Erythromycin (15)	77	76
Neomycin (40)	98	4
Tetracycline (100)	98	12

Adapted from Hagedorn et al. (1999)

Antibiotic Resistance Analysis (ARA)

Advantages

- Easy to type
- Easy to perform
- Easy to interpret
- Inexpensive

Disadvantages

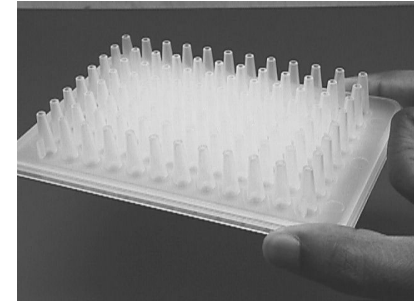
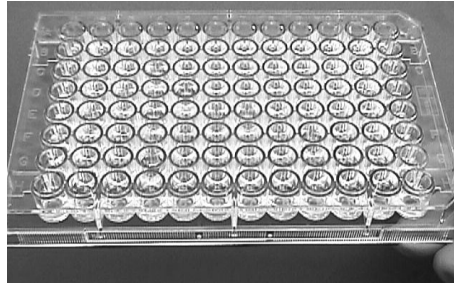
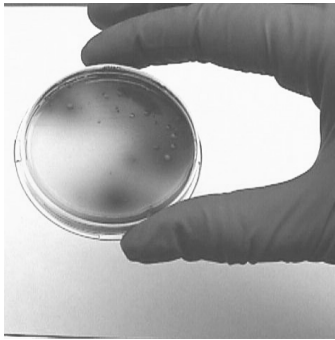
- Antibiotic resistance can be transferred
- Antibiotic dependent
- Culture dependent

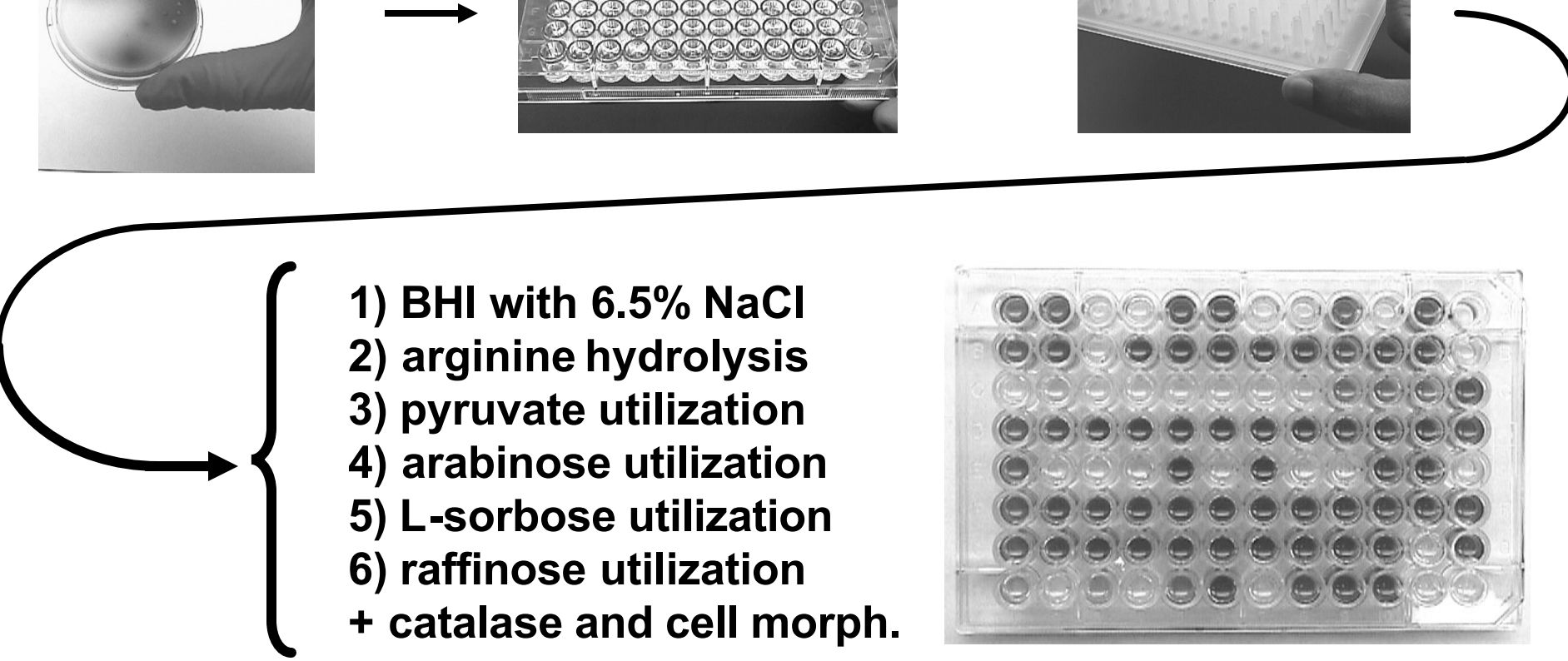
Carbon utilization (91 isolates + 5 controls)

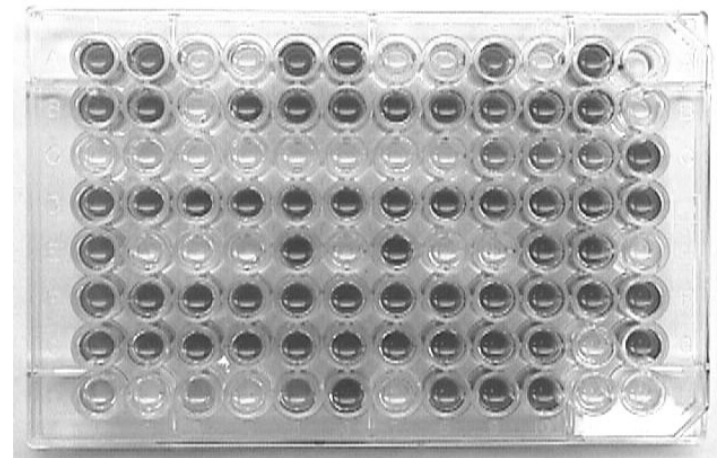
Enterococcosel agar

150 mL phosphate buffer

Replicator



- 
- 1) BHI with 6.5% NaCl
 - 2) arginine hydrolysis
 - 3) pyruvate utilization
 - 4) arabinose utilization
 - 5) L-sorbose utilization
 - 6) raffinose utilization
- + catalase and cell morph.



Carbon Utilization Profile

Advantages

- Easy to type
- Easy to perform
- Easy to interpret
- Relatively inexpensive
- Automated analysis

Disadvantages

- Unstable markers
- Ecologically irrelevant
- Culture dependent

Why molecular techniques?

- Genotyping methods are more sensitive and stable
- Do not rely on the culturability of microorganisms
- Simultaneous detection/monitoring of different groups without the requirement of a library

Steps in Fingerprinting Methods

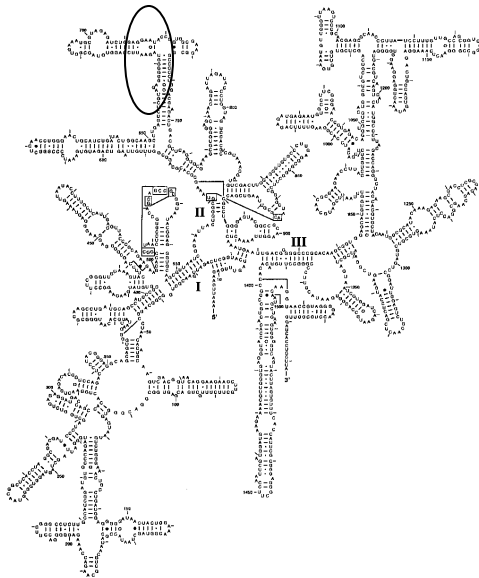
- Bacterial isolation
- ID confirmation
- DNA extractions
- PCR-amplification or DNA digestion
- Gel electrophoresis (and Southern hybridization)
- Image capture and analysis
- Average Rate of Correct Classification

DNA Fingerprinting Methods

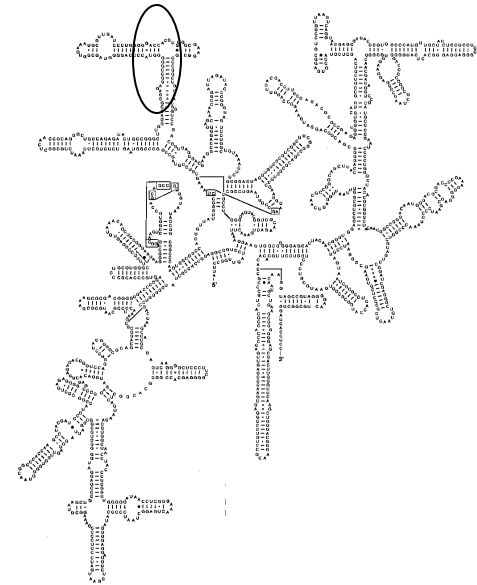
- RFLP analysis
- RAPD
- rep-PCR
- Sequencing ITS regions
- PCR-amplification of specific insertion sequences (IS)
- PFGE (gold standard in outbreak research)
- AFLP

Comparative Sequence Analysis of 16S rRNA

**16S rRNA of
*Escherichia coli***



**16S rRNA of
*Acinetobacter calcoaceticus***



	650	660	670
<i>E. coli</i>	GCUUGAGUCU	CGUAGAGGGG	GGUAGAAUUC
<i>A. calcoaceticus</i>	GCUAGAGUAU	GGGAGAGGAU	GGUAGAAUUC

MST Molecular Methods

Genotypic Methods – library dependent

- PFGE**
- Ribotyping**
- rep-PCR**
- AFLP**

MST Molecular Methods

- **Library independent methods**
 - **T-RFLP**
 - **LH-PCR**